


SEQUENCE LISTING

(1) GENERAL INFORMATION

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- (i) APPLICANT: ISNER, Jeffrey  
ASAHARA, Takayuki
  - (ii) TITLE OF THE INVENTION: METHODS FOR REGULATING ANGIOGENESIS  
(as amended)
  - (iii) NUMBER OF SEQUENCES: 2
  - (iv) CORRESPONDENCE ADDRESS:
    - (A) ADDRESSEE: DIKE, BRONSTEIN, ROBERTS & CUSMAN, LLP
    - (B) STREET: 130 Water Street
    - (C) CITY: Boston
    - (D) STATE: MA
    - (E) COUNTRY: USA
    - (F) ZIP: 02109
  - (v) COMPUTER READABLE FORM:
    - (A) MEDIUM TYPE: Diskette
    - (B) COMPUTER: IBM Compatible
    - (C) OPERATING SYSTEM: DOS
    - (D) SOFTWARE: FastSEQ for Windows Version 2.0
  - (vi) CURRENT APPLICATION DATA:
    - (A) APPLICATION NUMBER: 08/744,882
    - (B) FILING DATE: 08-NOV-1996
    - (C) CLASSIFICATION:
  - (vii) PRIOR APPLICATION DATA:
    - (A) APPLICATION NUMBER:
    - (B) FILING DATE:
  - (viii) ATTORNEY/AGENT INFORMATION:
    - (A) NAME: Resnick, David S
    - (B) REGISTRATION NUMBER: 34,235
    - (C) REFERENCE/DOCKET NUMBER: 46963
  - (ix) TELECOMMUNICATION INFORMATION:
    - (A) TELEPHONE: 617-523-3400
    - (B) TELEFAX: 617-523-6440
    - (C) TELEX:

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 30 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

AAGACATTTT CGGGCTCAG CTGCGCACCC

30

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TGGGGTAGGC ACTTTAGTAG TTCTCCTAAC

30

